

Figure 1

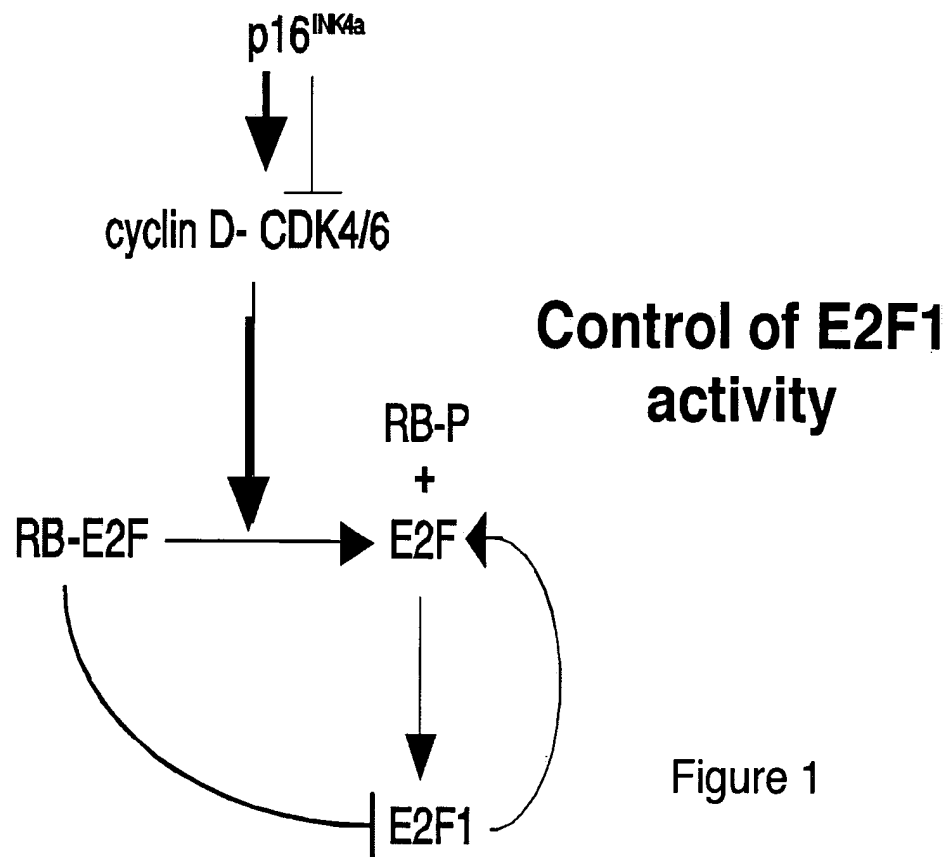


Figure 1

Figure 2A

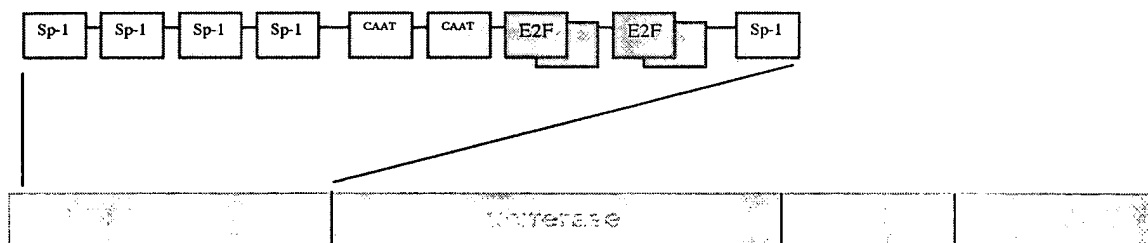


Figure 2B. E2F1 promoter sequence with EcoRI sites (273 bp w/o ER1). -208 to +66 from Johnson et al, Genes&Dev, 1994

GAATTCCATCCGGACAAAGCCTGCGCGCGCCCCGCCCCGCCATTGGCC
 GTACCGCCCCGCGCCGCGCCCCCATCCCGCCCCCTCGCCGCCGGGTCCG
 GCGCGTTAAAGCCAATAGGAACCGCCGCCGTTGTTCCCGTCACGGCC
 GGGGCAGCCAATTGTGGCGGCGCTCGGCGGCTCGTGGCTCTTTC**GCG**
GCAAAAAAGGATTTG**GCGCGTAAAA**GTGGCCGGGACTTTGCAGGCAGC
 GGCGGCCGGGGGCGGAGCGGGATCGAGCCCTCGCCGAGGCCTGAAT
 TC

Bases highlighted in yellow are primer sequences

C – missing base in construct

EcoR1

Stu1

Sp1 binding sites

CAAT box

Consensus E2F binding site (T/C)TT(C/G)(C/G)CG(C/G) (two imperfect palindromes)

G = +1 (transcription start)

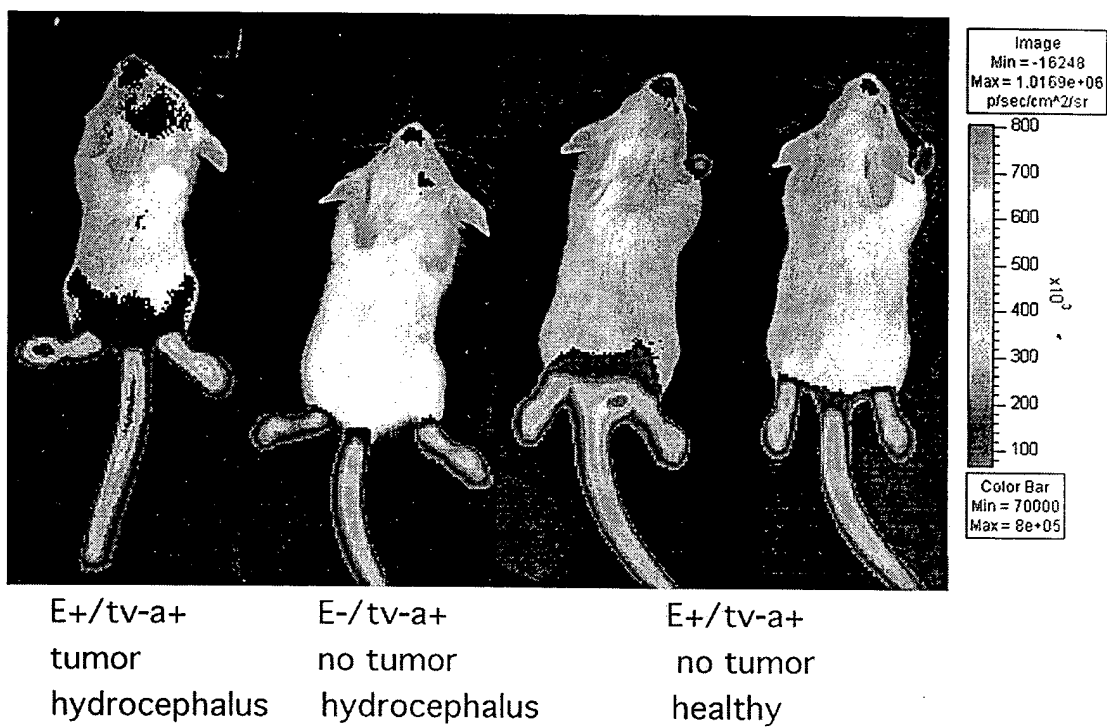


Fig. 3A

Figure 3B

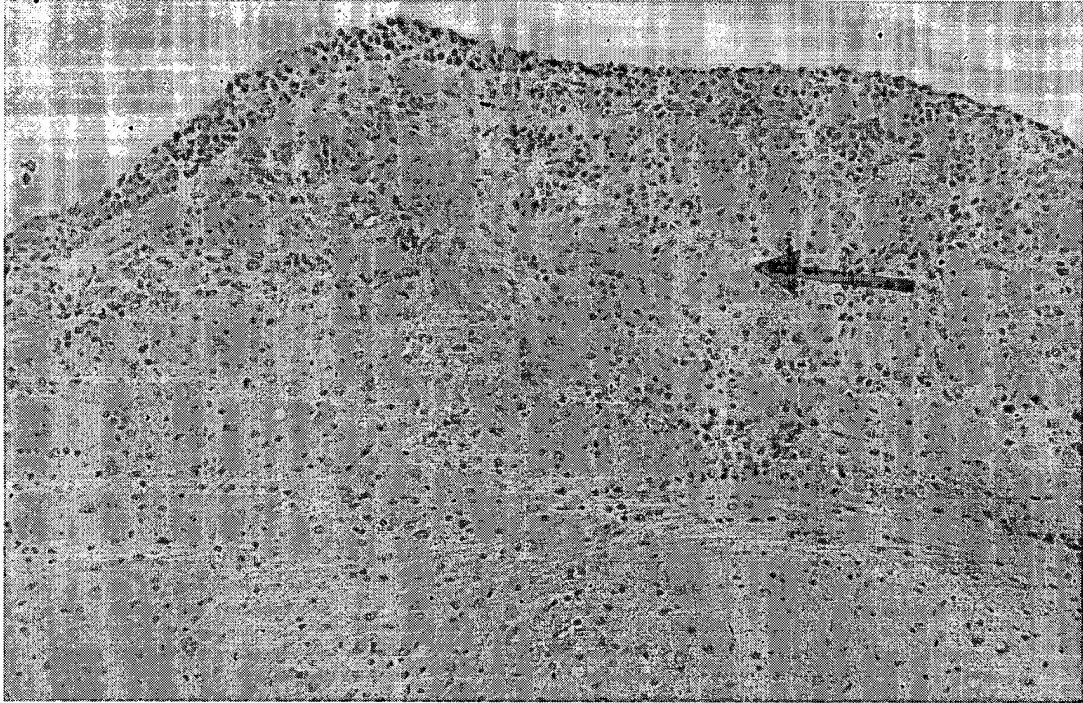
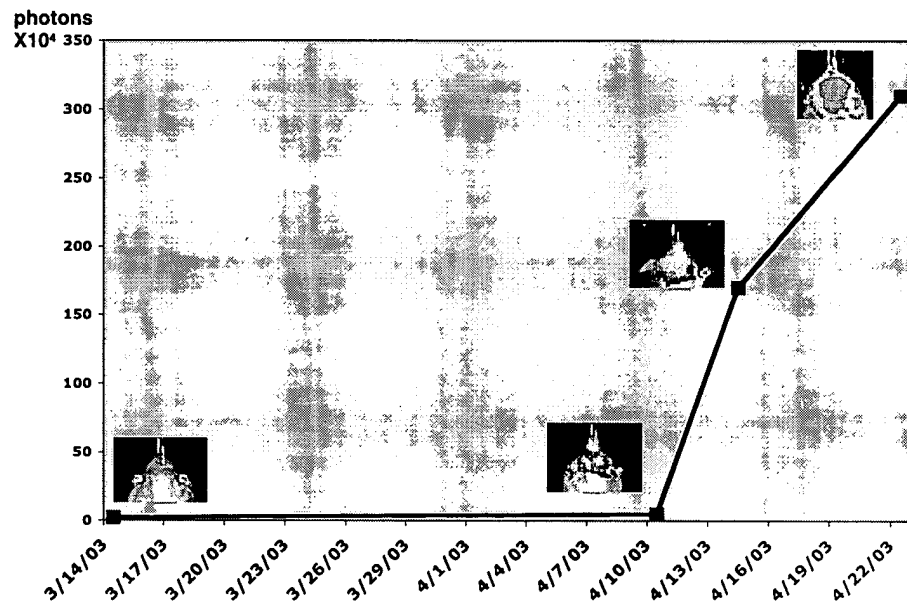


Figure 4



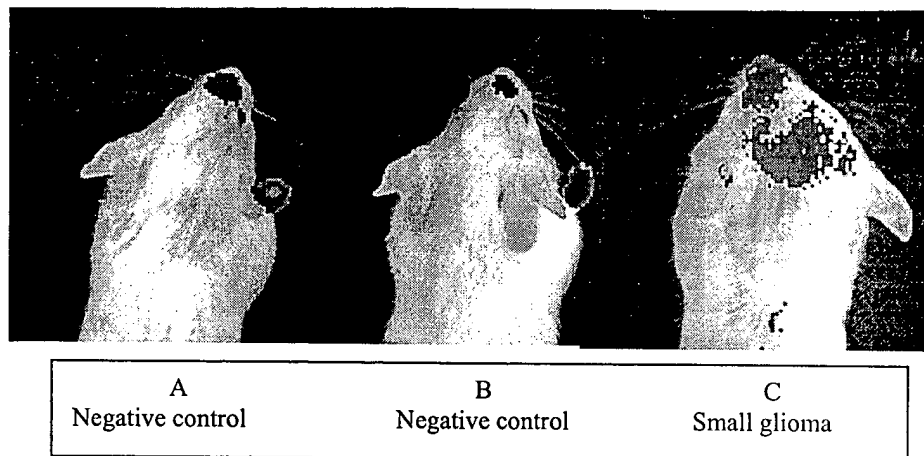


Fig. 5A

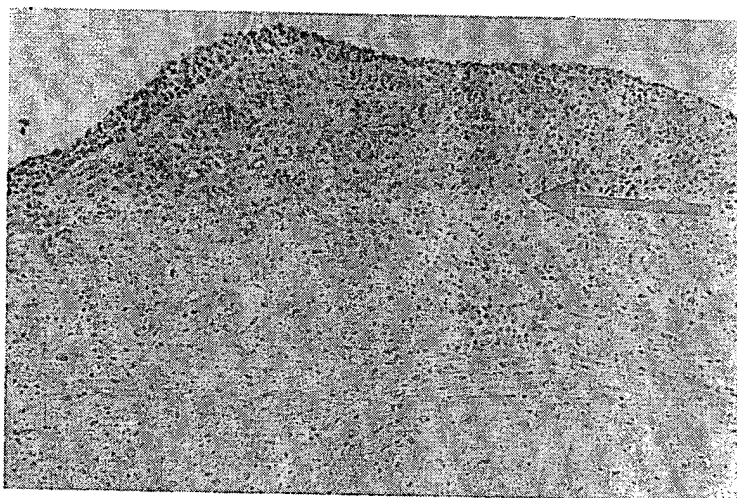


Fig. 5B

Figure 6

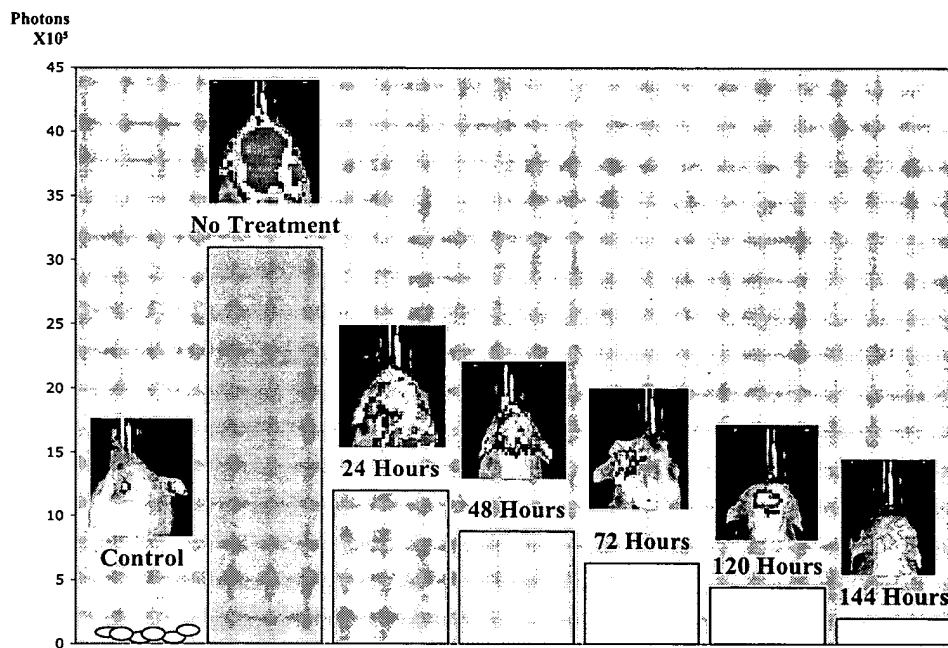


Figure 7A

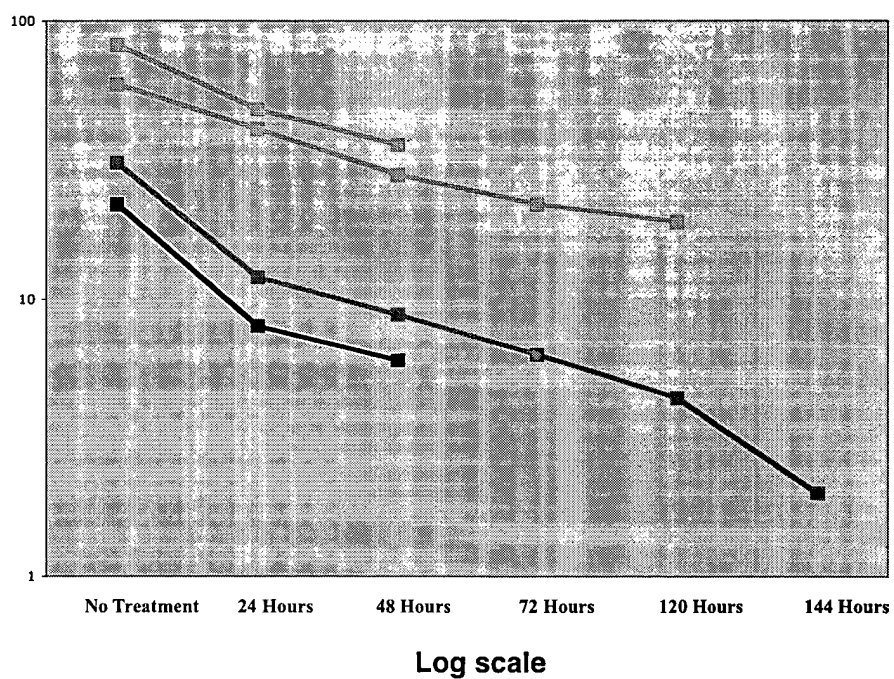


Figure 7B

